
Sequence Listing could not be accepted due to errors.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866)

217-9197 (toll free).

Reviewer: Durreshwar Anjum

Timestamp: [year=2009; month=1; day=30; hr=13; min=17; sec=42; ms=221;]

Reviewer Comments:

<150> JP 2004-122898

<151> 2004-04-19

<160> 75

The number provided for numeric identifier <160> must match the total number of sequences in the file. There were 75 sequences counted in this sequence listing. Numeric identifier <160> states there are a total of 74 sequences. Please make all necessary changes.

Validated By CRFValidator v 1.0.3

Application No: 10587431 Version No: 2.0

Input Set:

Output Set:

Started: 2009-01-14 21:17:26.254

Finished: 2009-01-14 21:17:36.647

Elapsed: 0 hr(s) 0 min(s) 10 sec(s) 393 ms

Total Warnings: 66

Total Errors: 29

No. of SeqIDs Defined: 75

Actual SeqID Count: 74

Error code		Error Description									
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(9)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(10)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(11)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(12)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(13)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(14)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(15)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(16)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(17)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(18)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(19)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(20)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(21)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(22)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(23)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(24)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(25)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(26)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(27)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(28)

Input Set:

Output Set:

Started: 2009-01-14 21:17:26.254 **Finished:** 2009-01-14 21:17:36.647

Elapsed: 0 hr(s) 0 min(s) 10 sec(s) 393 ms

Total Warnings: 66
Total Errors: 29
No. of SeqIDs Defined: 75

Actual SeqID Count: 74

Error code Error Description This error has occured more than 20 times, will not be displayed Ε 257 Invalid sequence data feature in <221> in SEQ ID (60) 257 Ε Invalid sequence data feature in <221> in SEQ ID (60) 257 Invalid sequence data feature in <221> in SEQ ID (60) Ε Ε 257 Invalid sequence data feature in <221> in SEQ ID (60) 257 Ε Invalid sequence data feature in <221> in SEQ ID (60) Ε 257 Invalid sequence data feature in <221> in SEQ ID (60) Ε 257 Invalid sequence data feature in <221> in SEQ ID (60) 257 Ε Invalid sequence data feature in <221> in SEQ ID (60) 257 Invalid sequence data feature in <221> Ε in SEQ ID (60) Invalid sequence data feature in <221> Ε 257 in SEQ ID (60) Ε 257 Invalid sequence data feature in <221> in SEQ ID (60) Ε 257 Invalid sequence data feature in <221> in SEQ ID (60) Ε 257 Invalid sequence data feature in <221> in SEQ ID (60) Ε 257 Invalid sequence data feature in <221> in SEQ ID (60) 257 Invalid sequence data feature in <221> Ε in SEQ ID (60) Ε 257 Invalid sequence data feature in <221> in SEQ ID (60) Ε 257 Invalid sequence data feature in <221> in SEQ ID (60) 257 Invalid sequence data feature in <221> Ε in SEQ ID (60) Ε 257 Invalid sequence data feature in <221> in SEQ ID (60) Ε 257 Invalid sequence data feature in <221> in SEQ ID (60) This error has occured more than 20 times, will not be displayed 252 \mathbf{E} Calc# of Seq. differs from actual; 75 seqIds defined; count=74

SEQUENCE LISTING

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<110> KAI, HIKARU
     TSUBAKI, MASAYUKI
      KUROKAWA, MASATO
<120> METHOD OF PRODUCING VIRUS
<130> 09864/0207778-US0
<140> 10587431
<141> 2009-01-14
<150> PCT/JP2005/007459
<151> 2005-04-19
<150> JP 2004-122898
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Pro Asp Ser Gly Arg
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<211> 7 <212> PRT

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Leu Gly Thr Ile Pro Gly
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<211> 10
<212> PRT
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Arg Asn Ile Ala Glu Ile Ile Lys Asp Ile
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Ile Lys Val Ala Val
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<213> Homo sapiens
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Asp Gly Glu Ala
<210> 9
<211> 10
<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic
      auxiliary amino acid sequence
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Gly Ala Gly Ala Gly Ala Gly Ala
1 5
               10
<210> 10
<211> 40
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
    auxiliary amino acid sequence
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                   10
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Gly Ala Gly Ala Gly Ala
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                 10
Gly Ala Gly Ala Gly Ala Gly Ala Gly Ala Gly Ala Gly Ala
                   25
        20
                                     30
Gly Ala Gly Ala Gly Ala Gly Ala Gly Ala Gly Ala Gly Ala
    35 40 45
Gly Ala Gly Ala
   50
             55
                                60
Gly Ala Gly Ala Gly Ala Gly Ala Gly Ala Gly Ala Gly Ala
65 70 75 80
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Gly Ala Gly Ala Gly Ala Gly Ala Gly Ala Gly Ala Gly Ala

90 95

Gly Ala Gly Al

Gly Ala 115 120 125

Gly Ala 130 135 140

Gly Ala 145 150 155 160

<210> 12

<211> 12

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic auxiliary amino acid sequence

<400> 12

Gly Ala Gly Ala Gly Ser Gly Ala Gly Ala Gly Ser

1 5 10

<210> 13

<211> 54

<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic auxiliary amino acid sequence

<400> 13

Gly Ala Gly Ala Gly Ser Gly Ala Gly Ala Gly Ser Gly Ala Gly Ala 1 5 10 15

Gly Ser Gly Ala Gly Ala Gly Ser Gly Ala Gly Ala Gly Ser Gly Ala
20 25 30

Gly Ala Gly Ser Gly Ala Gly Ala Gly Ser Gly Ala Gly Ala Gly Ser
35 40 45

Gly Ala Gly Ala Gly Ser

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<210> 14
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        20 25 30
Gly Ala Gly Ser Gly Ala Gly Ala Gly Ser Gly Ala Gly Ala Gly Ser
    35 40
                         45
Gly Ala Gly Ala Gly Ser Gly Ala Gly Ala Gly Ser Gly Ala Gly Ala
  50 55 60
Gly Ser Gly Ala Gly Ala Gly Ser Gly Ala Gly Ala Gly Ser Gly Ala
65 70 75 80
Gly Ala Gly Ser Gly Ala Gly Ala Gly Ser Gly Ala Gly Ala Gly Ser
         85 90 95
Gly Ala Gly Ala Gly Ser Gly Ala Gly Ala Gly Ser Gly Ala Gly Ala
       100 105 110
Gly Ser Gly Ala Gly Ala Gly Ser Gly Ala Gly Ala Gly Ser Gly Ala
            120
                          125
    115
Gly Ala Gly Ser Gly Ala Gly Ala Gly Ser Gly Ala Gly Ala Gly Ser
  130 135 140
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Gly Ala Gly Ala Gly Tyr Gly Ala Gly Ala Gly Tyr
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                                   10
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            20
                                25
Gly Ala Gly Tyr Gly Ala Gly Ala Gly Tyr Gly Ala Gly Ala Gly Tyr
        35
                            40
                                                45
Gly Ala Gly Ala Gly Tyr
   50
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10

Gly Tyr Gly Ala Gly Ala Gly Tyr Gly Ala Gly Ala Gly Tyr Gly A 20 25 30	la									
Gly Ala Gly Tyr Gly Ala Gly Ala Gly Tyr Gly Ala Gly Ala Gly T 35 40 45	yr									
Gly Ala Gly Ala Gly Tyr Gly Ala Gly Ala Gly Tyr Gly Ala Gly A 50 55 60	la									
Gly Tyr Gly Ala Gly Ala Gly Tyr Gly Ala Gly Ala Gly Tyr Gly A 65 70 75 8										
Gly Ala Gly Tyr Gly Ala Gly Ala Gly Tyr Gly Ala Gly Ala Gly T 85 90 95	yr									
Gly Ala Gly Ala Gly Tyr Gly Ala Gly Ala Gly Tyr Gly Ala Gly Al	la									
Gly Tyr Gly Ala Gly Ala Gly Tyr Gly Ala Gly Ala Gly Tyr Gly A 115 120 125	la									
Gly Ala Gly Tyr Gly Ala Gly Ala Gly Tyr Gly Ala Gly T 130 135 140	yr									
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1 5 10 15
Gly Tyr Gly Ala Gly Val Gly Tyr Gly Ala Gly Val Gly Tyr Gly Ala
       20
                25
Gly Val Gly Tyr Gly Ala Gly Val Gly Tyr Gly Ala Gly Val Gly Tyr
     35 40 45
Gly Ala Gly Val Gly Tyr
 50
<210> 20
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<212> PRT
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   5
                   10
Gly Tyr Gly Ala Gly Val Gly Tyr Gly Ala Gly Val Gly Tyr Gly Ala
      20 25 30
Gly Val Gly Tyr Gly Ala Gly Val Gly Tyr Gly Ala Gly Val Gly Tyr
                40
     35
                                    45
Gly Ala Gly Val Gly Tyr Gly Ala Gly Val Gly Tyr Gly Ala Gly Val
Gly Tyr Gly Ala Gly Val Gly Tyr Gly Ala Gly Val Gly Tyr Gly Ala
65
          70
                              75
```

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85 90 95
Gly Ala Gly Val Gly Tyr Gly Ala Gly Val Gly Tyr Gly Ala Gly Val
        100 105
                                        110
Gly Tyr Gly Ala Gly Val Gly Tyr Gly Ala Gly Val Gly Tyr Gly Ala
   115 120
Gly Val Gly Tyr Gly Ala Gly Val Gly Tyr Gly Ala Gly Val Gly Tyr
             135
Gly Ala Gly Val Gly Tyr Gly Ala Gly Val Gly Tyr Gly Ala Gly Val
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               150
                    155 160
Gly Tyr Gly Ala Gly Val Gly Tyr Gly Ala Gly Val Gly Tyr Gly Ala
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           165
                                          175
Gly Val Gly Tyr
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    auxiliary amino acid sequence
Gly Ala Gly Tyr Gly Val Gly Ala Gly Tyr Gly Val Gly Ala Gly Tyr
   5
                 10 15
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Gly Val Gly Ala Gly Tyr Gly Val Gly Ala Gly Tyr Gly Val Gly Ala

Gly Val Gly Tyr Gly Ala Gly Val Gly Tyr Gly Ala Gly Val Gly Tyr

20 25 30

Gly Tyr Gly Val Gly Ala Gly Tyr Gly Val Gly Ala Gly Tyr Gly Val 35 40 45

Gly Ala Gly Tyr Gly Val 50

<210> 23

<211> 180

<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic auxiliary amino acid sequence

<400> 23

Gly Ala Gly Tyr Gly Val Gly Ala Gly Tyr Gly Val Gly Ala Gly Tyr

1 10 15

Gly Val Gly Ala Gly Tyr Gly Val Gly Ala Gly Tyr Gly Val Gly Ala
20 25 30

Gly Tyr Gly Val Gly Ala Gly Tyr Gly Val Gly Ala Gly Tyr Gly Val 35 40 45

Gly Ala Gly Tyr Gly Val Gly Ala Gly Tyr Gly Val Gly Ala Gly Tyr
50 55 60

Gly Val Gly Ala Gly Tyr Gly Val Gly Ala Gly Tyr Gly Val Gly Ala 65 70 75 80

Gly Tyr Gly Val Gly Ala Gly Tyr Gly Val Gly Ala Gly Tyr Gly Val 85 90 95

Gly Ala Gly Tyr Gly Val Gly Ala Gly Tyr Gly Val Gly Ala Gly Tyr
100 105 110

Gly Val Gly Ala Gly Tyr Gly Val Gly Ala Gly Tyr Gly Val Gly Ala 115 120 125

Gly Tyr Gly Val Gly Ala Gly Tyr Gly Val Gly Ala Gly Tyr Gly Val 130 135 140

```
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145 150 155 160
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           165 170
Gly Tyr Gly Val
 180
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                10
      5
Ala Ala Ala Ala Gly Gly Ala Asp Gly Gly Ala Ala Ala Ala Ala
        20
            25
Ala Gly Gly Ala Asp Gly Gly Ala Ala Ala Ala Ala Gly Gly Ala
                                  45
    3.5
                  40
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<211> 18
<212> PRT
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Gly Ala
<210> 26
<211> 72
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<212> PRT

<213> Artificial Sequence

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<220>
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<223> Description of Artificial Sequence: Synthetic auxiliary amino acid sequence

<400> 26

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Ala Ala Ala Ala Gly Gly Ala 65 70

<210> 27

<211> 10

<212> PRT

<213> Artificial Sequence

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<400> 27

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1 5 10

<210> 28

<211> 50

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic auxiliary amino acid sequence

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Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro 35 40 45

Gly Val 50

<210> 29

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<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic auxiliary amino acid sequence

<400> 29

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1 5 10 15

Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val 20 25 30

Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro 35 40 45

Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly 50 55 60

Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val 65 70 75 80

Gly Val Pro Gly Val Gly Val Gly Val Pro Gly Val Gly
85 90 95

Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val 100 \$105\$

Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly V